

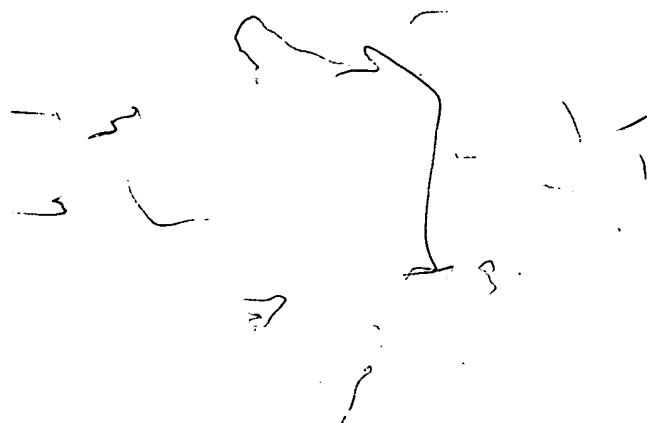
5040

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/517,956
Source: PCT
Date Processed by STIC: 12-28-04

ENTERED



RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/517,956

DATE: 12/28/2004

TIME: 11:41:47

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12282004\J517956.raw

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50 Ala Ala Lys Ile His Lys Lys Gly Met Ile Lys Ser Ser Arg Pro Leu
51 225          230          235          240
52 Arg Val Leu Thr Ala Val Val Ala Ser Phe Phe Ile Cys Trp Phe Pro
53          245          250          255
54 Phe Gln Leu Val Ala Leu Leu Gly Thr Val Trp Leu Lys Glu Met Leu
55          260          265          270
56 Phe Tyr Gly Lys Tyr Lys Ile Ile Asp Ile Leu Val Asn Pro Thr Ser
57          275          280          285
58 Ser Leu Ala Phe Phe Asn Ser Cys Leu Asn Pro Met Leu Tyr Val Phe
59          290          295          300
60 Val Gly Gln Asp Phe Arg Glu Arg Leu Ile His Ser Leu Pro Thr Ser
61 305          310          315          320
62 Leu Glu Arg Ala Leu Ser Glu Asp Ser Ala Pro Thr Asn Asp Thr Ala
63          325          330          335
64 Ala Asn Ser Ala Ser Pro Pro Ala Glu Thr Glu Leu Gln Ala Met
65          340          345          350
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68 <211> LENGTH: 1053
69 <212> TYPE: DNA
70 <213> ORGANISM: Human
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75 gtcctgggca atgggcttgt gatctgggtg gctggattcc ggatgacacg cacagtcacc 180
76 accatctgtt acctgaacct ggccctggct gacttttctt tcacggccac attaccattc 240
77 ctcatctgtt ccatggccat gggagaaaaa tggccttttg gctggttcct gtgtaagtta 300
78 attcacatcg tgggtggacat caacctcttt ggaagtgtct tcttgattgg ttctattgca 360
79 ctggaccgct gcatttgtgt cctgcatcca gtctgggccc agaaccaccg cactgtgagt 420
80 ctggccatga aggtgatcgt cggaccttgg attcttgctc tagtccttac cttgccagtt 480
81 ttctcttttt tgactacagt aactattcca aatgggggaca catactgtac tttcaacttt 540
82 gcatcctggg gtggcacccc tgaggagagg ctgaagggtg ccattaccat gctgacagcc 600
83 agagggatta tccggtttgt cattggcttt agcttgccga tgtccattgt tgccatctgc 660
84 tatgggctca ttgcagccaa gatccacaaa aagggcata ttaaataccag ccgtccctta 720
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87 gacatcctgg ttaacccaac gagctccctg gccttcttca acagctgcct caaccccatg 900
88 ctttacgtct ttgtgggcca agacttccga gagagactga tccactccct gccaccagt 960
89 ctggagaggg ccctgtctga ggactcagcc ccaactaatg acacggctgc caattctgct 1020
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93 <211> LENGTH: 24
94 <212> TYPE: PRT
95 <213> ORGANISM: Human
97 <400> SEQUENCE: 3
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99 1 5 10 15
100 Asp Leu Pro Val Lys Arg Arg Ala
101 20 24
103 <210> SEQ ID NO: 4

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104 <211> LENGTH: 24
105 <212> TYPE: PRT
106 <213> ORGANISM: Human
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111 Asp Leu Pro Val Lys Arg Arg Ala
112           20           24
114 <210> SEQ ID NO: 5
115 <211> LENGTH: 24
116 <212> TYPE: PRT
117 <213> ORGANISM: Human
119 <400> SEQUENCE: 5
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121   1           5           10           15
122 Asp Leu Pro Val Lys Arg Arg Thr
123           20
125 <210> SEQ ID NO: 6
126 <211> LENGTH: 21
127 <212> TYPE: PRT
128 <213> ORGANISM: Human
130 <400> SEQUENCE: 6
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134           20   21
136 <210> SEQ ID NO: 7
137 <211> LENGTH: 38
138 <212> TYPE: PRT
139 <213> ORGANISM: Rat
141 <400> SEQUENCE: 7
142 Met Ala Lys Arg Gly Phe Asn Cys Leu Leu Leu Ser Ile Ser Glu Ile
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145           20           25           30
146 Tyr Gly Ala Ser Ile Tyr
147           35           38
149 <210> SEQ ID NO: 8
150 <211> LENGTH: 24
151 <212> TYPE: PRT
152 <213> ORGANISM: Rat
154 <400> SEQUENCE: 8
155 Met Ala Lys Arg Gly Phe Asn Cys Leu Leu Leu Ser Ile Ser Glu Ile
156           5           10           15
157 Asp Leu Pro Val Lys Arg Leu Glu
158           20           24
160 <210> SEQ ID NO: 9
161 <211> LENGTH: 21
162 <212> TYPE: PRT

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163 <213> ORGANISM: Rat
165 <400> SEQUENCE: 9
166 Met Ala Lys Arg Gly Phe Asn Cys Leu Leu Leu Ser Ile Ser Glu Ile
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168 Asp Leu Pro Val Lys
169           20  21
171 <210> SEQ ID NO: 10
172 <211> LENGTH: 351
173 <212> TYPE: PRT
174 <213> ORGANISM: Rat
176 <400> SEQUENCE: 10
177 Met Glu Ala Asn Tyr Ser Ile Pro Leu Asn Val Ser Glu Val Val Val
178           5                      10                      15
179 Tyr Asp Ser Thr Ile Ser Arg Val Leu Trp Ile Leu Thr Met Val Val
180           20                      25                      30
181 Leu Ser Ile Thr Phe Val Leu Gly Val Leu Gly Asn Gly Leu Val Ile
182           35                      40                      45
183 Trp Val Ala Gly Phe Arg Met Val His Thr Val Thr Thr Thr Cys Phe
184           50                      55                      60
185 Leu Asn Leu Ala Leu Ala Asp Phe Ser Phe Thr Val Thr Leu Pro Phe
186           65                      70                      75                      80
187 Phe Val Ile Ser Ile Ala Met Lys Glu Lys Trp Pro Phe Gly Trp Phe
188           85                      90                      95
189 Leu Cys Lys Leu Val His Ile Val Val Asp Ile Asn Leu Phe Gly Ser
190           100                     105                     110
191 Val Phe Leu Ile Ala Leu Ile Ala Leu Asp Arg Cys Ile Cys Val Leu
192           115                     120                     125
193 His Pro Val Trp Ala Gln Asn His Arg Thr Val Ser Leu Ala Arg Lys
194           130                     135                     140
195 Val Val Val Gly Pro Trp Ile Leu Ala Leu Ile Leu Thr Leu Pro Ile
196           145                     150                     155                     160
197 Phe Ile Phe Met Thr Thr Val Arg Ile Pro Gly Gly Asn Val Tyr Cys
198           165                     170                     175
199 Thr Phe Asn Phe Ala Ser Trp Gly Asn Thr Ala Glu Glu Leu Leu Asn
200           180                     185                     190
201 Ile Ala Asn Thr Phe Val Thr Val Arg Gly Ser Ile Arg Phe Ile Ile
202           195                     200                     205
203 Gly Phe Ile Met Pro Met Ser Ile Val Ala Ile Cys Tyr Gly Leu Ile
204           210                     215                     220
205 Ala Val Lys Ile His Arg Arg Ala Leu Val Asn Ser Ser Arg Pro Leu
206           225                     230                     235                     240
207 Arg Val Leu Thr Ala Val Val Ala Ser Phe Phe Ile Cys Trp Phe Pro
208           245                     250                     255
209 Phe Gln Leu Val Ala Leu Leu Gly Thr Ile Trp Phe Lys Glu Ser Leu
210           260                     265                     270
211 Phe Ser Gly Arg Tyr Lys Ile Leu Asp Met Trp Val His Pro Thr Ser
212           275                     280                     285
213 Ser Leu Ala Tyr Phe Asn Ser Cys Leu Asn Pro Met Leu Tyr Ala Phe
214           290                     295                     300

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215 Met Gly Gln Asp Phe His Glu Arg Leu Ile His Ser Leu Pro Ser Ser
216 305                      310                      315                      320
217 Leu Glu Arg Ala Leu Ser Glu Asp Ser Gly Gln Thr Ser Asp Thr Gly
218                      325                      330                      335
219 Ile Ser Ser Ala Leu Pro Pro Val Asn Ile Asp Ile Lys Ala Ile
220                      340                      345                      350
222 <210> SEQ ID NO: 11
223 <211> LENGTH: 1053
224 <212> TYPE: DNA
225 <213> ORGANISM: Rat
227 <400> SEQUENCE: 11
228 atggaagcca actattccat ccctctgaat gtatcagaag tgggtgtcta tgattctacc 60
229 atctccagag ttttgtggat cctcacaatg gtggttctct ccatcacctt tgtcctgggt 120
230 gtgctgggta atggactagt gatctgggta gctggattcc ggatgggtaca cactgtcacc 180
231 actacctgtt ttctgaatct agctttggct gacttctctt tcacagtgac tctaccattc 240
232 tttgtcatct caattgctat gaaagaaaaa tggccttttg gatggttcct gtgtaaatta 300
233 gttcacattg tagtagacat aaacctcttt ggaagtgtct tcctgattgc tttaattgcc 360
234 ttggaccgct gcatttgtgt cctgcatcca gtctgggctc agaaccaccg cactgtgagc 420
235 ctggctagga aggtggttgt tgggccctgg attttagctc tgattctcac tttgcccatt 480
236 tttattttca tgactacagt tagaattcct ggaggcaatg tgtactgtac attcaacttc 540
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238 agaggagca tcaggttcat tattggcttc ataatgccta tgtccattgt tgccatctgc 660
239 tatggactca tcgctgtcaa gatccacaga agagcacttg ttaattccag ccgtccatta 720
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241 gcccttttag gtacaatctg gtttaaagag tcattgttta gtggtcgtta caaaattctt 840
242 gacatgtggg ttcacccaac cagctcattg gcctacttca atagttgcct caatccaatg 900
243 ctctatgctt tcatgggcca ggactttcat gaaagactga ttcattccct gccttccagt 960
244 ctggagagag ccctgagtga ggactctggc caaaccagtg atacaggcat cagttctgct 1020
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248 <211> LENGTH: 351
249 <212> TYPE: PRT
250 <213> ORGANISM: Mouse
252 <400> SEQUENCE: 12
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255 Tyr Asp Ser Thr Ile Ser Arg Val Leu Trp Ile Leu Ser Met Val Val
256                      20                      25                      30
257 Val Ser Ile Thr Phe Phe Leu Gly Val Leu Gly Asn Gly Leu Val Ile
258                      35                      40                      45
259 Trp Val Ala Gly Phe Arg Met Pro His Thr Val Thr Thr Ile Trp Tyr
260                      50                      55                      60
261 Leu Asn Leu Ala Leu Ala Asp Phe Ser Phe Thr Ala Thr Leu Pro Phe
262 65                      70                      75                      80
263 Leu Leu Val Glu Met Ala Met Lys Glu Lys Trp Pro Phe Gly Trp Phe
264                      85                      90                      95
265 Leu Cys Lys Leu Val His Ile Val Val Asp Val Asn Leu Phe Gly Ser
266                      100                      105                      110
267 Val Phe Leu Ile Ala Leu Ile Ala Leu Asp Arg Cys Ile Cys Val Leu

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/517,956

DATE: 12/28/2004

TIME: 11:41:48

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12282004\J517956.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application No

L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date